

Final TCR Display PCT.WorkFile.txt

Organization Applicant

Street : 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : martin.green@avidex.com
<110> OrganizationName : Avidex Ltd

Individual Applicant

Street : Avidex Ltd, 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : bent.jakobsen@avidex.com
<110> LastName : Jakobsen
<110> FirstName : Bent
<110> MiddleInitial : K
<110> Suffix : Dr

Individual Applicant

Street : Avidex Ltd, 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : torben.andersen@avidex.com
<110> LastName : Andersen
<110> FirstName : Torben
<110> MiddleInitial : B
<110> Suffix : Dr

Individual Applicant

Street : Avidex Ltd, 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : peter.molloy@avidex.com
<110> LastName : Molloy
<110> FirstName : Peter
<110> MiddleInitial : E
<110> Suffix : Dr

Individual Applicant

Street : Avidex Ltd, 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : jonathan.boulter@avidex.com
<110> LastName : Boulter

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<110> FirstName : Jonathan
<110> MiddleInitial : M
<110> Suffix : Dr

Individual Applicant

Street : Avidex Ltd, 57C Milton Park
City : Abingdon
State : Oxfordshire
Country : England
PostalCode : OX14 4RX
PhoneNumber : +44 1235 438600
FaxNumber : +44 1235 438601
EmailAddress : yi.li@avidex.com

<110> LastName : Li
<110> FirstName : Yi
<110> MiddleInitial :
<110> Suffix : Dr

Application Project

<120> Title : Substances
<130> AppFileReference : Case No 19
<140> CurrentAppNumber : GB 0226227.7
<141> CurrentFilingDate : 2002-11-09

Earlier Applications

<150> PriorAppNumber : GB0226227.7
<151> PriorFilingDate : 2002-11-09

Earlier Applications

<150> PriorAppNumber : GB 0301814.0
<151> PriorFilingDate : 2003-01-25

Earlier Applications

<150> PriorAppNumber : GB 0304067.2
<151> PriorFilingDate : 2003-02-22

Earlier Applications

<150> PriorAppNumber : GB 0311397.4
<151> PriorFilingDate : 2003-05-16

Earlier Applications

<150> PriorAppNumber : GB 0316356.5
<151> PriorFilingDate : 2003-07-11

Earlier Applications

<150> PriorAppNumber : US 60/436,323
<151> PriorFilingDate : 2003-04-16

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DSDVYITDKT VLDMRSMDFK
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 1
SequenceDescription :

Sequence

20

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
QSKDSDVYIT DKTVLDMRSM
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 2
    SequenceDescription :

Sequence
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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DIQNPDPAVY QLRDSKSSDK
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 3
    SequenceDescription :

Sequence
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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DPAVYQLRDS KSSDKSVCLF
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 4
    SequenceDescription :

Sequence
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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
NGKEVHSGVS TDPQPLKEQP
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 5
    SequenceDescription :

Sequence
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ALNDSRYALS SRLRVSATFW
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 6
    SequenceDescription :

Sequence
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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PPEVAVFEPS EAEISHTQKA
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 7
    SequenceDescription :

Sequence
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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KEVHSGVSTD PQPLKEQPAL
<212> Type : PRT
<211> Length : 20
    SequenceName : SEQ ID 8
    SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
VFPPEVAVFE PSEAEISHTQ 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 9
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PYIQNPEAV YQLKDPRSQD STLCLFTDFD SQINVPKTME SGTFITDKTV LDMKAMDSKS 60
NGAIAWSNQT SFTCQDIFKE TNATYPSSDV P 91
<212> Type : PRT
<211> Length : 91
SequenceName : SEQ ID 10
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
EDLRNVTPPK VSLFEPSKAE IANKQKATLV CLARGFFPDH VELSWWVNNGR EVHSGVSTD 60
P QAYKESNYSY CLSSRLRVSA TFWHNPRNHF RCQVQFHGLS EEDKWPEGSP KPVTQNISAE 120
AWGRAD 126
<212> Type : PRT
<211> Length : 126
SequenceName : SEQ ID 11
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ESGTFITDKT VLDMKAMDSK 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 12
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KTMESGTFIT DKTVLDMKAM 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 13
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
YIQNPEAVY QLKDPRSQDS 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 14
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVYQLKDPRS QDSTLCLFTD 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 15
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
NGREVHSGVS TDPQAYKESN 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 16
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KESNYSYCLS SRLRVSATFW 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 17
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PPKVSLFEPS KAEIANKQKA 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 18
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
REVHSGVSTD PQAYKESNYS 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 19
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
VTPPKVSLFE PSKAEIANKQ 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 20
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LLFGYPVYV 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 21
SequenceDescription :

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Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
SLLMITQC
<212> Type : PRT
<211> Length : 8
SequenceName : SEQ ID 22
SequenceDescription :

8

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LLGRNSFEV
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 23
SequenceDescription :

9

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KLVALGINAV
<212> Type : PRT
<211> Length : 10
SequenceName : SEQ ID 24
SequenceDescription :

10

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LLGDLFGV
<212> Type : PRT
<211> Length : 8
SequenceName : SEQ ID 25
SequenceDescription :

8

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
FLRGRAYGL
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 26
SequenceDescription :

9

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
HRCQAIRKK
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 27
SequenceDescription :

9

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
YRSGIIAVV
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 28

9

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SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
VYGFVRACL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 29
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ILAKFLHWL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 30
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LTLGEFLKL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 31
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PKYVKQNTLK LA 12
<212> Type : PRT
<211> Length : 12
SequenceName : SEQ ID 32
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
GILGFVFTL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 33
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
SLYNTVATL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 34
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cacagacaaa tgtgtgctag acat 24
<212> Type : DNA

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<211> Length : 24
SequenceName : SEQ ID 35
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 35

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgtctagca cacatttgc tgtg
<212> Type : DNA
<211> Length : 24
SequenceName : SEQ ID 36
SequenceDescription :

24

Custom Codon

Sequence Name : SEQ ID 36

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagtgggtc tgcacagacc c
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 37
SequenceDescription :

21

Custom Codon

Sequence Name : SEQ ID 37

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gggtctgtgc agacccact g
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 38
SequenceDescription :

21

Custom Codon

Sequence Name : SEQ ID 38

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
taataatacg tataataata ttctatttca aggagacagt c
<212> Type : DNA
<211> Length : 41
SequenceName : SEQ ID 39
SequenceDescription :

41

Custom Codon

Sequence Name : SEQ ID 39

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

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caatccagcg gctgccgtag gcaataggta tttcattatg actgtctcct tgaaatag 58

<212> Type : DNA
<211> Length : 58
SequenceName : SEQ ID 40
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 40

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctacggcagc cgctggattg ttattactcg cggcccatggcc gccatggcc cag 53

<212> Type : DNA
<211> Length : 53
SequenceName : SEQ ID 41
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 41

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gttctgctcc acttccttct gggccatggc cggctggcc g 41
<212> Type : DNA
<211> Length : 41
SequenceName : SEQ ID 42
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 42

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagaaggaag tggagcagaa c 21
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 43
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 43

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cttcttaaag aattcttaat taaccttaggt tattaggaac tttctggct gggaaag 57

<212> Type : DNA
<211> Length : 57
SequenceName : SEQ ID 44
SequenceDescription :

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Custom Codon

Sequence Name : SEQ ID 44

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :
gttaattaag aattcttaa gaaggagata tacatatgaa aaaatttatta ttgcatttc 60

<212> Type : DNA

<211> Length : 60

SequenceName : SEQ ID 45

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 45

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :
cgcgtgtga gaatagaaag gaacaactaa aggaattgcg aataataatt ttttcatatg 60

<212> Type : DNA

<211> Length : 60

SequenceName : SEQ ID 46

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 46

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :
ctttctattc tcacagcgcg caggctggtg tcactcagac 40

<212> Type : DNA

<211> Length : 40

SequenceName : SEQ ID 47

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 47

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :
atgatgtcta gatgcggccg cgtctgctct accccaggcc tc 42

<212> Type : DNA

<211> Length : 42

SequenceName : SEQ ID 48

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 48

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

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gcatctagac atcatcacca tcatactactag actgttgaaa gttgttagc aaaac 55

<212> Type : DNA
<211> Length : 55
SequenceName : SEQ ID 49
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 49

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctagaggta ccttattaa actccttatt acgcagttatg 40
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 50
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 50

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
agctgcagct aatacgactc actataggaa caggccacca tggcgatcgat tatgctgagt 60
gatattccttg tccgggttta cccttag 86
<212> Type : DNA
<211> Length : 86
SequenceName : SEQ ID 51
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 51

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gatccccatgg tggcctgttc ctatagttagtgc tcgtattttgc tgc 43
<212> Type : DNA
<211> Length : 43
SequenceName : SEQ ID 52
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 52

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
agctgcagct aatacgactc actataggaa caggccacca tgg 43
<212> Type : DNA
<211> Length : 43
SequenceName : SEQ ID 53
SequenceDescription :

Custom Codon

Final TCR Display PCT.WorkFile.txt
Sequence Name : SEQ ID 53

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ccaccatggg ccagaaggaa gtggaggcaga actc
<212> Type : DNA
<211> Length : 34
SequenceName : SEQ ID 54
SequenceDescription :

34

Custom Codon

Sequence Name : SEQ ID 54

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cgagagcccg tagaactgga cttg
<212> Type : DNA
<211> Length : 24
SequenceName : SEQ ID 55
SequenceDescription :

24

Custom Codon

Sequence Name : SEQ ID 55

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gtggatccgg cggtggcggg tcgaacgctg gtgtcactca gacccc
<212> Type : DNA
<211> Length : 46
SequenceName : SEQ ID 56
SequenceDescription :

46

Custom Codon

Sequence Name : SEQ ID 56

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ccgatccac ctccgcctga accgcctcca ccggtagcca caacctgggt ccctg
<212> Type : DNA
<211> Length : 55
SequenceName : SEQ ID 57
SequenceDescription :

55

Custom Codon

Sequence Name : SEQ ID 57

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctgagaattc ttatgactct ccgcgggttga agctc
<212> Type : DNA
<211> Length : 35
SequenceName : SEQ ID 58

35

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 58

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgacgaattc tgactctccg cggttgaagc tc
<212> Type : DNA
<211> Length : 32
SequenceName : SEQ ID 59
SequenceDescription :

32

Custom Codon

Sequence Name : SEQ ID 59

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
agctgcagct aatacgactc actatagg
<212> Type : DNA
<211> Length : 28
SequenceName : SEQ ID 60
SequenceDescription :

28

Custom Codon

Sequence Name : SEQ ID 60

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ggccaccatg ggcaacgctg gtgtcactca gacccc
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 61
SequenceDescription :

36

Custom Codon

Sequence Name : SEQ ID 61

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgaaccgcct ccaccgtctg ctctacccca ggcctcggcg
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 62
SequenceDescription :

40

Custom Codon

Sequence Name : SEQ ID 62

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgactctccg cggttgaagc tc
<212> Type : DNA

22

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<211> Length : 22
<400> PreSequenceString :
SequenceName : SEQ ID 63
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 63

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagctggggg aagttcagt ttggagcag
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 64
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 64

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctgctccaaa ctgaagcttc cccagctg
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 65
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 65

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gtacttctgt gcctcgaggc cgggactag
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 66
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 66

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctagtcccggt cctcgaggca cagaagtac
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 67
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 67

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
29

29

29

29

29

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catttcagg gatagcaagg
<212> Type : DNA
<211> Length : 20
SequenceName : SEQ ID 68
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 68

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tcacacagga aacagctatg tcacacagga aacagctatg 40
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 69
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 69

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggnnnnnknn knnnnnnnk cgaccagagc agtacttcg 49
<212> Type : DNA
<211> Length : 49
SequenceName : SEQ ID 70
SequenceDescription :

Feature

Sequence: SEQ ID 70:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 49
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 70

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccgnnnknn knnnnnnnk nnkccagagc agtacttcgg gc 52
<212> Type : DNA
<211> Length : 52
SequenceName : SEQ ID 71
SequenceDescription :

Feature

Sequence: SEQ ID 71:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

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Feature

Sequence: SEQ ID 71:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 52
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 71

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccgnnknn knnknnknnk nnkcgaccag agcagtactt cg 52
<212> Type : DNA
<211> Length : 52
SequenceName : SEQ ID 72
SequenceDescription :

Feature

Sequence: SEQ ID 72:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 52
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 72

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccgnnknn knnknnknnk nnkggaggc gaccagagca g 51
<212> Type : DNA
<211> Length : 51
SequenceName : SEQ ID 73
SequenceDescription :

Feature

Sequence: SEQ ID 73:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 51
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 73

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccgggann knnknnknnk nnknnkggc gaccagagca gtac 54

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```
<212> Type : DNA
<211> Length : 54
    SequenceName : SEQ ID 74
    SequenceDescription :

Feature
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Sequence: SEQ ID 74:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
    Other Information : m is a or c; n is a or t or g or c; k is g or t
    CDSJoin : No

Feature
-----
Sequence: SEQ ID 74:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 54
    Other Information : m is a or c; n is a or t or g or c; k is g or t
    CDSJoin : No

Custom Codon
-----
Sequence Name : SEQ ID 74

Sequence
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggnnknnknn knnnknnknnk ccagagcagt acttcgggc          49
<212> Type : DNA
<211> Length : 49
    SequenceName : SEQ ID 75
    SequenceDescription :

Feature
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Sequence: SEQ ID 75:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
    Other Information : m is a or c; n is a or t or g or c; k is g or t
    CDSJoin : No

Feature
-----
Sequence: SEQ ID 75:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 49
    Other Information : m is a or c; n is a or t or g or c; k is g or t
    CDSJoin : No

Custom Codon
-----
Sequence Name : SEQ ID 75

Sequence
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggnnknnknn knnnknnknnk gagcagtact tcgggccg          48
<212> Type : DNA
<211> Length : 48
```

Final TCR Display PCT.WorkFile.txt

SequenceName : SEQ ID 76

SequenceDescription :

Feature

Sequence: SEQ ID 76:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 48

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 76

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgtgcctcga ggnnnnnnn knnnnnnnn cagtacttcg ggccgggc 48

<212> Type : DNA

<211> Length : 48

SequenceName : SEQ ID 77

SequenceDescription :

Feature

Sequence: SEQ ID 77:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 48

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 77

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgtgcctcga ggccgnnnnn knnnnnnkggg cgaccagagc agtacttcg 49

<212> Type : DNA

<211> Length : 49

SequenceName : SEQ ID 78

SequenceDescription :

Feature

Sequence: SEQ ID 78:

<221> FeatureKey : misc_feature

<222> LocationFrom :

<222> LocationTo :

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Feature

Sequence: SEQ ID 78:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 49

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Final TCR Display PCT.WorkFile.txt

Sequence Name : SEQ ID 78

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmmnnnnnnm nmnnnnnntgt aacggcacag agtag 46
<212> Type : DNA
<211> Length : 46
SequenceName : SEQ ID 79
SequenceDescription :

Feature

Sequence: SEQ ID 79:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 46
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 79

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmmnnnnngc tgtcmnnntgt aacggcacag agtag 46
<212> Type : DNA
<211> Length : 46
SequenceName : SEQ ID 80
SequenceDescription :

Feature

Sequence: SEQ ID 80:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 46
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 80

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmmnnnnnnm ngctgtcmnn tgtaacggca cagaggtag 49
<212> Type : DNA
<211> Length : 49
SequenceName : SEQ ID 81
SequenceDescription :

Feature

Sequence: SEQ ID 81:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Final TCR Display PCT.WorkFile.txt

Feature

Sequence: SEQ ID 81:

<221> FeatureKey : misc_feature
 <222> LocationFrom : 1
 <222> LocationTo : 49
 Other Information : m is a or c; n is a or t or g or c; k is g or t
 CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 81

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 aaactgaagc ttmmnnnnngc tgcmmnnaac ggcacagagg tag 43
 <212> Type : DNA
 <211> Length : 43
 SequenceName : SEQ ID 82
 SequenceDescription :

Feature

Sequence: SEQ ID 82:

<221> FeatureKey : misc_feature
 <222> LocationFrom : 1
 <222> LocationTo : 43
 Other Information : m is a or c; n is a or t or g or c; k is g or t
 CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 82

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cgggactagc gggagggcga ccagagcagt ag 42
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 83
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 83

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 ASRPGLAGGR PEQY 14
 <212> Type : PRT
 <211> Length : 14
 SequenceName : SEQ ID 84
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cggggctgat gagtgcgtag ccagagcagt ac 42
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 85

Final TCR Display PCT.workFile.txt

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 85

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLMSAX PEQY 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 86
SequenceDescription :

Feature

Sequence: SEQ ID 86:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggctgag gtcggcgtag ccagagcagt ac 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 87
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 87

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLRSAX PEQY 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 88
SequenceDescription :

Feature

Sequence: SEQ ID 88:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggacttagc gggagggcga ccagaggcgt ag 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 89

Final TCR Display PCT.WorkFile.txt

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 89

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PEAX
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 90
SequenceDescription :

14

Feature

Sequence: SEQ ID 90:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Feature

Sequence: SEQ ID 90:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggactagc gggagggcga ccagaggatt ag
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 91
SequenceDescription :

42

Custom Codon

Sequence Name : SEQ ID 91

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PEDX
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 92
SequenceDescription :

14

Feature

Sequence: SEQ ID 92:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon

Final TCR Display PCT.WorkFile.txt

CDSJoin : No

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cgggactagc gggagggcga ccagatcagt ag
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 93
 SequenceDescription :

42

Custom Codon

 Sequence Name : SEQ ID 93

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 ASRPGLAGGR PDQX
 <212> Type : PRT
 <211> Length : 14
 SequenceName : SEQ ID 94
 SequenceDescription :

14

Feature

 Sequence: SEQ ID 94:
 <221> FeatureKey : MISC_FEATURE
 <222> LocationFrom :
 <222> LocationTo :
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Feature

 Sequence: SEQ ID 94:
 <221> FeatureKey : MISC_FEATURE
 <222> LocationFrom : 1
 <222> LocationTo : 14
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cgggtctgtt ggctgggcga ccagagcagt ac
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 95
 SequenceDescription :

42

Custom Codon

 Sequence Name : SEQ ID 95

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 ASRPGLXAGR PEQY
 <212> Type : PRT
 <211> Length : 14
 SequenceName : SEQ ID 96

14

Final TCR Display PCT.WorkFile.txt
SequenceDescription :

Feature

Sequence: SEQ ID 96:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Feature

Sequence: SEQ ID 96:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggctggc tccggggcga ccagagcagt ag 42
<212> Type : DNA
<211> Length : 42
 SequenceName : SEQ ID 97
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 97

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLVPGR PEQX 14
<212> Type : PRT
<211> Length : 14
 SequenceName : SEQ ID 98
 SequenceDescription :

Feature

Sequence: SEQ ID 98:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Feature

Sequence: SEQ ID 98:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
 Other Information : x denotes an amino acid residue encoded by an amber codon
 CDSJoin : No

Sequence

Final TCR Display PCT.WorkFile.txt

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggcttgt gtctgcttag ccagagcagt ac 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 99
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 99

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLVSAK PEQY 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 100
SequenceDescription :

Feature

Sequence: SEQ ID 100:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggacttagc gggagggcga ccacatccgt ag 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 101
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 101

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PHPX 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 102
SequenceDescription :

Feature

Sequence: SEQ ID 102:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Feature

Final TCR Display PCT.WorkFile.txt

Sequence: SEQ ID 102:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggactagc gggagggcga ccagatgcgt ag
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 103
SequenceDescription :

42

Custom Codon

Sequence Name : SEQ ID 103

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PDAX
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ 104
SequenceDescription :

14

Feature

Sequence: SEQ 104:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggtctgat tagtgcttag ccagagcagt ac
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 105
SequenceDescription :

42

Custom Codon

Sequence Name : SEQ ID 105

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLISAX PEQY
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 106
SequenceDescription :

14

Feature

Final TCR Display PCT.WorkFile.txt

Sequence: SEQ ID 106:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggaaagctt cag 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 107
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 107

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL Q 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 108
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggccgctt cag 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 109
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 109

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL Q 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 110
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggaaagatg cag 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 111
SequenceDescription :

Custom Codon

Final TCR Display PCT.WorkFile.txt
Sequence Name : SEQ ID 111

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKM Q 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 112
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggaaagttg cat 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 113
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 113

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL H 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 114
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg ggggttagctt cat 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 115
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 115

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGXL H 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 116
SequenceDescription :

Feature

Sequence: SEQ ID 116:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber codon

Final TCR Display PCT.WorkFile.txt

CDSJoin : No

Feature

Sequence: SEQ ID 116:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 11
Other Information : x denotes an amino acid residue encoded by an amber codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggggagctt cat 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 117
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 117

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGEL H 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 118
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg ggggaggctg cat 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 119
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 119

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGRL H 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 120
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg ggggcagctt cat 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 121

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 121

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGQL H
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 122
SequenceDescription :

11

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggaaagggtt cat
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 123
SequenceDescription :

33

Custom Codon

Sequence Name : SEQ ID 123

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKV H
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 124
SequenceDescription :

11

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccgttacaa ctgacagctg gggaaagggtg aat
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 125
SequenceDescription :

33

Custom Codon

Sequence Name : SEQ ID 125

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKV N
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 126
SequenceDescription :

11

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

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gccgttacaa ctgacagctg gggaaagctt ctg 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 127
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 127

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL L 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 128
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ggaattcatc gatgcagaag gaagtggagc ag 32
<212> Type : DNA
<211> Length : 32
SequenceName : SEQ ID 129
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 129

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gtacacggcc gggtcagggt tctggatata c 31
<212> Type : DNA
<211> Length : 31
SequenceName : SEQ ID 130
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 130

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tctctcatta atgaatgctg gtgtcactca gacccc 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 131
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 131

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tagaaaaccgg tggccaggca caccagtgtg gc 32
<212> Type : DNA

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<211> Length : 32
SequenceName : SEQ ID 132
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 132

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctgcctctggt tccgcactc
<212> Type : DNA
<211> Length : 19
SequenceName : SEQ ID 133
SequenceDescription :

19

Custom Codon

Sequence Name : SEQ ID 133

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gagtgccgaa ccagagcag
<212> Type : DNA
<211> Length : 19
SequenceName : SEQ ID 134
SequenceDescription :

19

Custom Codon

Sequence Name : SEQ ID 134

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atatccagaa cccggatcct gccgtgta
<212> Type : DNA
<211> Length : 28
SequenceName : SEQ ID 135
SequenceDescription :

28

Custom Codon

Sequence Name : SEQ ID 135

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tacacacggcag gaatccgggt tctggatat
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 136
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 136

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

Final TCR Display PCT.WorkFile.txt

ggagatatac atatgcagga ggtgacacag	30
<212> Type : DNA	
<211> Length : 30	
SequenceName : SEQ ID 137	
SequenceDescription :	
Custom Codon	

Sequence Name : SEQ ID 137	
Sequence	

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
tacacggcag gatccgggtt ctggatatt	29
<212> Type : DNA	
<211> Length : 29	
SequenceName : SEQ ID 138	
SequenceDescription :	
Custom Codon	

Sequence Name : SEQ ID 138	
Sequence	

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
ggagatatac atatgggtgt cactcagacc	30
<212> Type : DNA	
<211> Length : 30	
SequenceName : SEQ ID 139	
SequenceDescription :	
Custom Codon	

Sequence Name : SEQ ID 139	
Sequence	

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
cccaagctta gtctgctcta ccccaggcct cgcc	34
<212> Type : DNA	
<211> Length : 34	
SequenceName : SEQ ID 140	
SequenceDescription :	
Custom Codon	

Sequence Name : SEQ ID 140	
Sequence	

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
gccggccatg gccaaacagg aggtgacgca gattcct	37
<212> Type : DNA	
<211> Length : 37	
SequenceName : SEQ ID 141	
SequenceDescription :	
Custom Codon	

Sequence Name : SEQ ID 141	
Sequence	

Final TCR Display PCT.WorkFile.txt

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cttcttaaag aattcttaat taaccttaggt tattaggaac tttctgggct ggggaag 57

<212> Type : DNA
<211> Length : 57
SequenceName : SEQ ID 142
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 142

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tcacagcgcg caggctggtg tcactcagac cccaaa 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 143
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 143

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cgagagcccg tagaactgga ct tg 24
<212> Type : DNA
<211> Length : 24
SequenceName : SEQ ID 144
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 144

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tctctcatta atgaatgctg gtgtcactca gacccc 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 145
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 145

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tagaaaccgg tggccaggca caccagtgtg g 31
<212> Type : DNA
<211> Length : 31
SequenceName : SEQ ID 146
SequenceDescription :

Custom Codon

Final TCR Display PCT.WorkFile.txt

Sequence Name : SEQ ID 146

Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgcagaagg aagtggagca gaactctgga cccctcagtg ttccagaggg agccattgcc      60
tctctcaact gcacttacag tgaccgaggt tcccagtcct tcttctggta cagacaatat      120
tctggaaaaa gccctgagtt gataatgtcc atatactcca atggtgacaa agaagatgga      180
aggtttacag cacagctcaa taaagccagc cagtatgtt ctctgctcat cagagactcc      240
cagcccagtg attcagccac ctacctctgt gccgttacaa ctgacagctg gggaaattg      300
cagtttggag cagggaccca ggttgtggtc accccagata tccagaaccc tgaccctgccc 360
gtgtaccagc tgagagactc taaatccagt gacaagtctg tctgcctatt caccgatttt      420
gattctcaaa caaatgtgtc acaaagtaag gattctgtatg tgtatatcac agacaaatgt      480
gtgctagaca tgaggtctat ggacttcaag agcaacagtg ctgtggcctg gagcaacaaa 540
tctgactttg catgtgcaaa cgccctcaac aacagcatta ttccagaaga cacccttctc 600
cccagcccag aaagttccta a                                              621
<212> Type : DNA
<211> Length : 621
      SequenceName : SEQ ID 147
      SequenceDescription :

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Custom Codon

Sequence Name : SEQ ID 147

Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
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gtccccatg gctacaatgt ctccagatca accacagagg atttcccgct caggctgctg      240
tcggctgctc cctcccagac atctgtgtac ttctgtgcca gcaggccggg actagcggga      300
gggcgaccag agcagtactt cgggcccggc accaggctca cggtcacaga ggacctgaaa      360
aacgtgttcc cacccgaggt cgctgtgtt gagccatcag aagcagagat ctcccacacc      420
caaaaggcca cactgggtg cctggccaca ggcttctacc ccgaccacgt ggagctgagc      480
tggtgtgtga atgggaagga ggtgcacagt ggggtctgca cagacccgca gcccctcaag      540
gagcagcccg ccctcaatga ctccagatac gctctgagca gccgcctgag ggtctcgccc 600
accttctggc aggacccccc caaccacttc cgctgtcaag tccagttcta cgggctctcg      660
gagaatgacg agtggaccca ggatagggcc aaaccgtca cccagatcgt cagcgccgag      720
gcctgggtta gaggcagacta a                                              741
<212> Type : DNA
<211> Length : 741
      SequenceName : SEQ ID 148

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Final TCR Display PCT.WorkFile.txt
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 148

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

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RFTAQLNKAS QYVSLIIRDS QPSDSATYLC AVTTDSWGKL QFGAGTQVVV TPDIQNPDPA 120

VYQLRDSKSS DKSVCFLFTDF DSQTNVSQSK DSDVYITDKC VLDMRSMDFK SNSAVAWSNK 180

SDFACANAFN NSIIIPEDTFF PSPESS 206

<212> Type : PRT

<211> Length : 206

SequenceName : SEQ ID 149

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

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NVFPPEVAVF EPSEAEISHT QKATLVCLAT GFYPDHVELS WWVNGKEVHS GVCTDPQPLK 180

EQPALNDSRY ALSSRLRVSA TFWQDPRNHF RCQVQFYGLS ENDEWTQDRA KPVTQIVSAE 240

AWGRAD

<212> Type : PRT

<211> Length : 246

SequenceName : SEQ ID 150

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

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ataaatattga aaaaggaaga gtatgagtat tcaacatttc cgtgtcgccc ttattccctt 180

ttttcggca tttgccttc ctgttttgc tcacccagaa acgctggtga aagtaaaaga 240

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ggggatcat gtaactcgcc ttgatcggtt ggaaccggag ctgaatgaag ccataccaaa 660

cgacgagcgt gacaccacga tgcctgttagc aatggcaaca acgttgcgc aactattaac 720

Final TCR Display PCT.WorkFile.txt

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Final TCR Display PCT.WorkFile.txt

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gcgg	ttctg	gggtggcgg	tctgagg	gcgg	ttccgg	4800

Final TCR Display PCT.WorkFile.txt

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<212> Type : DNA
<211> Length : 5495
SequenceName : SEQ ID 151
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 151

Sequence

Final TCR Display PCT.WorkFile.txt

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aatgacgagt	ggacccagga	tagggccaaa	cccgtcaccc	agatcgttag	cgccgaggcc	1140
tgggttagag	cagacggtgg	aggcggttca	ctcagcagca	ccctgacgct	gagcaaagca	1200
gactacgaga	aacacaaagt	ctacgcctgc	gaagtcaccc	atcagggcct	gagttcgccc	1260
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<212> Type : DNA
 <211> Length : 1303
 SequenceName : SEQ ID 152
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 152

Sequence

<213> OrganismName : Synthetic Sequence	
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SGGGGSNAGV TQTPKFQVLK TGQSMTLQCA QDMNHEYMSW YRQDPGMGLR LIHYSVGAGI	180
TDQGEVPNGY NVSRSTTEDF PLRLLSAAPS QTSVYFCASR PGLAGGRPEQ YFGPGTRLTV	240
TEDLKNVFPP EVAVFEPSEA EISHTQKATL VCLATGFYPD HVELSWWVNG KEVHSGVSTD	300
PQPLKEQPAL NDSRYALSSR LRVSATFWQD PRNHFRCQVQ FYGLSENDEW TQDRAKPVTQ	360
IVSAEAWGRA DGGGGSLSST LTLSKADYEK HKVYACEVTH QGLSSPVTKS FNRGES	416

<212> Type : PRT
 <211> Length : 416
 SequenceName : SEQ ID 153
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
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aatctgcag tccaccgtga aaagccctt tacacgcgcc ttgggataa acaaataaaa	120
agatttatgt aagtttatac ataggcgagt actctgttat tggactatt tacgaagtta	180
ttataacttt ttccctctca tactcataag ttgtaaaggc acagcggaa taagggaaaa	240
aacgcgtaa aacggaagga caaaaacgag tgggtcttgc cgaccactt catttctac	300
gacttctagt caacccacgt gctcacccaa tggtagttga ccttagagttg tcgccattct	360
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tgataagagt cttactgaac caactcatga gtggtcagtg tctttcgta gaatgcctac	540
cgtactgtca ttctcttaat acgtcacgac ggtattggta ctcactattg tgacgccggt	600

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Final TCR Display PCT.WorkFile.txt
 ctggcagagg ccctcgacgt acacagtctc caaaaagtggc agtagtggct ttgcgcgct 2699

<212> Type : DNA
 <211> Length : 2699
 SequenceName : SEQ ID 154
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 154

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
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 aatattctgg gaaaagccct gagttgataa tgtccatata ctccaatggg gacaaagaag 180
 atggaaagggt tacagcacag ctcaataaaag ccagccagta tggttctctg ctcatcagag 240
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 atttcccgct caggctgctg tcggctgctc cctcccgacatctgtgtac ttctgtgcca 660
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 ccgaccacgt ggagctgagc tggtggtgta atggaaagga ggtgcacagt ggggtcagca 900
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 gcccgcctgag ggtctcgcc accctctggc aggaccccg caaccacttc cgctgtcaag 1020
 tccagttcta cgggctctcg gagaatgacg agtggaccca ggatagggcc aaaccgtca 1080
 cccagatcgt cagcgccgag gcctgggtta gagcagacgg tggaggcgggt tcactcagca 1140
 gcaccctgac gctgagcaaa gcagactacg agaaacacaa agtctacgac tgcgaagtca 1200
 cccatcaggg cctgagttcg cccgtcacaa agagcttcaa ccgcggagag tcataagaat 1260
 tc 1262
 <212> Type : DNA
 <211> Length : 1262
 SequenceName : SEQ ID 155
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 155

Final TCR Display PCT.WorkFile.txt

Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg      60
cagtgtgccc agatatgaa ccatgaatac atgtcctggc atcgacaaga cccaggcatg      120
gggctgaggc tgattcatta ctcagttggc gctggatca ctgaccaagg agaagtcccc      180
aatggctaca atgtctccag atcaaccaca gaggattcc cgctcaggct gctgtcggct      240
gctccctccc agacatctgt gtacttctgt gccagcaggc cgggactagc gggagggtga      300
ccagagcagt acttcgggccc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg      360
ttcccacccg aggtcgctgt gttttagcca tcagaagcag agatctccca cacccaaaag      420
gccacactgg tgtgcctggc cacaggcttc tacccgacc acgtggagct gagctgggtgg      480
gtgaatggga aggaggtgca cagtgggtc tgacagacc cgcagccct caaggagcag      540
cccgccctca atgactccag atacgctctg agcagccgccc tgagggtctc ggccaccttc      600
tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat      660
gacgagtggc cccaggatag ggccaaacccc gtcacccaga tcgtcagcgc cgaggcctgg      720
ggtagagcag ac                                         732
<212> Type : DNA
<211> Length : 732
      SequenceName : SEQ ID 156
      SequenceDescription :
```

Custom Codon

```
-----
Sequence Name : SEQ ID 156
```

Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AGVTQTPKFQ VLKTGQSMAL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP      60
NGYNVSRSTT EDFPLRLLSA APSQTSVYFC ASRPGLAGGX PEQYFGPGTR LTVTEDLKNV      120
FPPEVAVFEP SAEISHTQK ATLVLATGF YPDHVELSWW VNGKEVHSGV CTDPQPLKEQ      180
PALNDSRYAL SSRLRVSATF WQDPRNHFRC QVQFYGLSEN DEWTQDRAKP VTQIVSAEAW      240
GRAD                                         244
<212> Type : PRT
<211> Length : 244
      SequenceName : SEQ ID 157
      SequenceDescription :
```

Feature

```
-----
Sequence: SEQ ID 157:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 244
      Other Information : X denotes an amino acid residue encoded by an opal
      codon, this is normally translated as a tryptophan
      CDSJoin : No
```

Sequence

Final TCR Display PCT.WorkFile.txt

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg 60
 tagtgtgccc aggatatgaa ccatgaatac atgtccttgtt atcgacaaga cccaggcatg 120
 gggctgagggc tgattcatta ctcagtttgtt gctggtatca ctgaccaagg agaagtcccc 180
 aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcaggct gctgtcggct 240
 gctccctccc agacatctgt gtacttctgt gccagcaggc cgggacttagc gggagggcga 300
 ccagagcagt acttcgggccc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg 360
 ttcccacccg aggtcgctgt gttttagcca tcagaagcag agatctccca cacccaaaag 420
 gcccacactgg tgtgcctggc cacaggcttc taccggacc acgtggagct gagctggtgg 480
 gtgaatggga aggaggtgca cagtgggttc tgcacagacc cgccagccct caaggagcag 540
 cccgcccctca atgactccag atacgctctg agcagccgccc tgagggtctc ggccacccctc 600
 tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat 660
 gacgagtggaa cccaggatag ggccaaacccc gtcacccaga tcgtcagcgc cgaggcctgg 720
 ggttagagcag ac 732
 <212> Type : DNA
 <211> Length : 732
 SequenceName : SEQ ID 158
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 158

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg 60
 cagtgtgccc aggatatgaa ccatgaatac atgtccttgtt atcgacaaga cccaggcatg 120
 gggctgagggc tgattcatta ctcagtttgtt gctggtatca ctgaccaagg agaagtcccc 180
 aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcaggct gctgtcggct 240
 gctccctccc agacatctgt gtacttctgt gcctcgaggc cgggctgtat gagtgcggaa 300
 ccagagcagt acttcgggccc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg 360
 ttcccacccg aggtcgctgt gttttagcca tcagaagcag agatctccca cacccaaaag 420
 gcccacactgg tgtgcctggc caccggtttc taccggacc acgtggagct gagctggtgg 480
 gtgaatggga aggaggtgca cagtgggttc tgcacagacc cgccagccct caaggagcag 540
 cccgcccctca atgactccag atacgctctg agcagccgccc tgagggtctc ggccacccctc 600
 tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat 660
 gacgagtggaa cccaggatag ggccaaacccc gtcacccaga tcgtcagcgc cgaggcctgg 720
 ggttagagcag actaagcttg aattc 745
 <212> Type : DNA
 <211> Length : 745
 SequenceName : SEQ ID 159

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 159

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

MNAGVTQTPK FQVLKTGQSM TLQCAQDMNH EYMSWYRQDP GMGLRLIHYS VGAGITDQGE	60
VPNGYNVSR S TTEDFPLRLL SAAPSQTTSVY FCASRPGLMS AEPEQYFGPG TRLTVTEDLK	120
NVFPPEVAVF EPSEAEISHT QKATLVCLAT GFYPDHVELS WWVNGKEVHS GVCTDPQPLK	180
EQPALNDSRY ALSSRLRVSA TFWQDPRNHF RCQVQFYGLS ENDEWTQDRA KPVTQIVSAE	240

AWGRAD

<212> Type : PRT

<211> Length : 246

SequenceName : SEQ ID 160

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AGVTQTPKFQ VLKTGQSMTL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP	60
NGYNVSRSTT EDFPLRLLSA APSQTTSVYFC ASRPGLMSAQ PEQYFGPGTR LTVTEDLKNV	120
FPPEVAVFEP SEAEISHTQK ATLVLATGF YPDHVELSWW VNGKEVHSGV CTDPQPLKEQ	180
PALNDSRYAL SSRLRVSATF WQDPRNHFRC QVQFYGLSEN DEWTQDRAKP VTQIVSAEAW	240

GRAD

<212> Type : PRT

<211> Length : 244

SequenceName : SEQ ID 161

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

atgcaggagg tgacacagat tcctgcagct ctgagtgtcc cagaaggaga aaacttggtt	60
ctcaactgca gtttcactga tagcgctatt tacaacctcc agtggtttag gcaggaccct	120
ggaaagggtc tcacatctct gttgcttatt cagtcaagtc agagagagca aacaagtgg	180
agacttaatg cctcgctgga taaatcatca ggacgttagta ctttatacat tgcagcttct	240
cagcctggtg actcagccac ctacctctgt gctgtgaggc ccacatcagg aggaagctac	300
atacctacat ttggaagagg aaccagcctt attgttcatc cgtatatcca gaaccctgac	360
cctgccgtgt accagctgag agactctaaa tccagtgaca agtctgtctg cctattcacc	420
gattttgatt ctcaaacaaa tgtgtcacaa agtaaggatt ctgatgtgta tattcacagac	480
aaatgtgtgc tagacatgag gtctatggac ttcaagagca acagtgtgt ggcctggagc	540
aacaatctg actttgcatt tgcaaacgcc ttcaacaaca gcattattcc agaagacacc	600
ttcttccccca gcccagaag ttccctaa	627

<212> Type : DNA

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<211> Length : 627
 SequenceName : SEQ ID 162
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 162

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 atgggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg 60
 cagtgtgccc agatatgaa ccatgaatac atgtcctggc atcgacaaga cccaggcatg 120
 gggctgaggg tgattcatta ctcagttggc gctggtatca ctgaccaagg agaagtcccc 180
 aatggctaca atgtctccag atcaaccaca gaggattcc cgctcaggct gctgtccggct 240
 gctccctccc agacatctgt gtacttctgt gccagcagtt acgtcgggaa caccggggag 300
 ctgtttttg gagaaggctc taggctgacc gtactggagg acctgaaaaa cgtgttccca 360
 cccgaggtcg ctgtgttga gccatcagaa gcagagatct cccacaccca aaaggccaca 420
 ctggtgtgcc tggccacagg cttctacccc gaccacgtgg agctgagctg gtgggtgaat 480
 gggaggagg tgcacagtgg ggtctgcaca gacccgcagc ccctcaagga gcagcccgcc 540
 ctcaatgact ccagatacgc tctgagcagc cgccctgaggg tctcggccac cttctggcag 600
 gaccccgca accacttccg ctgtcaagtc cagttctacg ggctctcgga gaatgacgag 660
 tggaccagg atagggccaa acccgtcacc cagatcgtca gcgcggaggc ctgggtttaga 720
 gcagactaa 729

<212> Type : DNA
 <211> Length : 729
 SequenceName : SEQ ID 163
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 163

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 MQEVTVQIPAA LSVPEGENLV LNCSFTDSAI YNLQWFRQDP GKGLTSLLI QSSQREQTSG 60
 RLNASLDKSS GRSTLYIAAS QPGDSATYLC AVRPTSGGSY IPTFGRGTSV IVHPYIQNPD 120
 PAVYQLRDSK SSDKSVCCLFT DFDSQTNVSQ SKDSDVYITD KCVLDMRSMD FKSNSAVAWS 180
 NKSDFACANA FNNSIIIPEDT FFPSPESS 208

<212> Type : PRT
 <211> Length : 208
 SequenceName : SEQ ID 164
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 MGVTQTPKFQ VLKTGQSMTL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP 60

Final TCR Display PCT.WorkFile.txt

NGYNVSRSTT	EDFPLRLLSA	APSQTSVYFC	ASSYVGNTGE	LFVGEGSRLT	VLEDLKNVFP	120
PEVAVFEPSE	AEISHTQKAT	LVCLATGFYP	DHVELSWWVN	GKEVHSGVCT	DPQPLKEQPA	180
LNDSRYALSS	RLRVSATFWQ	DPRNHFRCQV	QFYGLSENDE	WTQDRAKPVT	QIVSAEAWGR	240
AD						242
<212> Type : PRT						
<211> Length : 242						
SequenceName : SEQ ID 165						
SequenceDescription :						
Sequence						

<213> OrganismName : Synthetic Sequence						
<400> PreSequenceString :						
ttcctggcct	tttgctggcc	tttgctcac	atgtaatgtg	agtttagctca	aaggaccgga	60
aaacgaccgg	aaaacgagtg	tacattacac	tcaatcgagt	ctcattaggc	accccaggct	120
ttacacttta	tgcttccggc	tcgtatgtt	gagtaatccg	tggggtccga	aatgtgaaat	180
acgaaggccg	agcataacaac	tgtggattt	tgagcggata	acaatttac	acaggaaaca	240
gctatgacca	acaccttaac	actcgccat	tgttaaagt	tgtcctttgt	cgatactgg	300
tgattacgcc	aagctacgta	cttaagtatt	ctatttcaag	gagacagtca	actaatgcgg	360
ttcgatgcat	gaattcataa	gataaagttc	ctctgtcagt	taatgaaata	cctattgcct	420
acggcagccg	ctggattgtt	attactcgcg	attactttat	ggataacgga	tgccgtcggc	480
gacctaacaa	taatgagcgc	gcccagccgg	ccatggccaa	acaggaggtg	acgcagattc	540
ctgcagctct	cgggtcggcc	ggtaccgggt	tgtcctccac	tgcgtctaag	gacgtcgaga	600
gagtgccca	gaaggagaaa	acttggttct	caactgcagt	ttcactgata	ctcacagggt	660
cttcctcttt	tgaaccaaga	gttgacgtca	aagtgactat	gcgctattt	caacctccag	720
tggtttaggc	aggaccctgg	gaaaggcttc	cgcgataaat	gttggaggtc	accaaattccg	780
tcctgggacc	ctttccagag	acatctctgt	tgcttattca	gtcaagtcag	agagagcaaa	840
caagtggaaag	tgttagagaca	acgaataagt	cagttcagtc	tctctcgttt	gttcacccccc	900
acttaatgcc	tcgctggata	aatcatcagg	acgtagttact	ttatacattt	tgaattacgg	960
agcgacctat	ttagtagtcc	tgcatacatga	aatatgtaac	cagttctca	gcctgggtgac	1020
tcagccacct	acctctgtgc	tgtgaggccc	gtcgaagagt	cggaccactg	agtcgggtgga	1080
tggagacacg	acactccggg	acatcaggag	gaagctacat	acctacattt	ggaagaggaa	1140
ccagccttat	tgttagtcctc	cttcgatgt	tggatgtaaa	ccttctcattt	ggtcggaaata	1200
tgttcatccg	tatatccaga	acccggatcc	tgccgtgtac	cagctgagag	acaagttaggc	1260
atataggct	tgggcctagg	acggcacatg	gtcgactctc	actctaaatc	cagtgacaag	1320
tctgtctgcc	tattcaccga	ttttgattct	tgagatttag	gtcactgttc	agacagacgg	1380
ataagtggct	aaaactaaga	caaacaatg	tgtcacaag	taaggattct	gatgtgtata	1440
tcacagacaa	gtttgtttac	acagtgtttc	attcctaaga	ctacacatat	agtgtctgtt	1500
atgtgtgcta	gacatgaggt	ctatggactt	caagagcaac	agtgcgtgg	tacacacgat	1560

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ctgtactcca	gatacctgaa	gttctcggt	tcacgacacc	cctggagcaa	caaatctgac	1620	
tttgcatgt	caaacgcctt	caacaacagc	ggacctcg	tttagactg	aaacgtacac	1680	
gttgcggaa	gttgcgtcg	attattccag	aagacacctt	cttccccagc	ccagaaagtt	1740	
cctaataacc	taataagg	tc ttgtggaa	gaagggtcg	gttcttcaa	ggattattgg	1800	
taggttaatt	aagaattctt	taagaagggg	atatacatat	aaaaaaat	atccaattaa	1860	
ttcttaagaa	attcttccc	tatatgtata	ctttttat	ttattcgca	ttcccttagt	1920	
tgttccccc	tattctcaca	gcgcgcaggc	aataagcgtt	aaggaaatca	acaaggaaag	1980	
ataagagtgt	cgcgcgtccg	tgggtcact	cagacccaa	aattccaggt	cctgaagaca	2040	
ggacagagca	accacagtga	gtctgggtt	ttaagg	gttcca	ggacttctgt	2100	
tgacactgca	gtgtgc	ccag	gatatgaacc	atgaatacat	gtcctggat	actgtgacgt	2160
cacacgggtc	ctataacttgg	tacttatgt	caggaccata	cgacaagacc	caggcatgg	2220	
gctgaggctg	attcattact	cagttgg	tc	gcttctgg	gtccgtaccc	cgactccgac	2280
taagtaatga	gtcaaccacg	tgg	tatcact	gaccaaggag	aagtcccaa	tggctacaat	2340
gtctccagat	accatagtga	ctgg	ttcc	ttcagg	gggttacc	accgtgtt	2400
caaccacaga	ggat	ttcc	cg	ctcagg	ctgc	tgcggctgc	2460
cctaaagg	gagtccgacg	acagccgacg	agggagg	gttgg	gtgtct	acatctgtgc	2520
cagcagttac	gtcgg	aaaca	ccggg	gagct	tgt	tgacacacg	2580
cagcccttgc	ggccc	ctcg	gtt	tttgg	gaa	aggctcta	2640
ctgaaaaacg	caaaa	ac	ttcc	ggagat	ccgact	ttcgac	2700
tgttcccacc	cgagg	tcg	gtt	tttg	act	ggaggac	2760
gctccagcga	caca	aaact	gt	actt	tttgc	actggagg	2820
ggtgtcctg	gcc	acagg	tct	cccc	gt	ttgggtt	2880
cggtgtccga	agat	gggg	cc	acgtgg	gt	ggtaatgg	2940
cacagtgggg	ggtgcac	ctc	gact	cgacca	cc	acttacc	3000
tctgcacaga	ccc	gcag	cc	tcagg	tt	ccac	3060
ggcg	tcgg	gg	gtt	actg	gg	tgagc	3120
cctgagg	tcgg	ccac	tct	ggcagg	tct	gagcc	3180
agccgg	ttgg	cc	cact	ccgt	gt	caagtcc	3240
ctctcg	gggg	ccgtt	gt	gaagg	tt	caagatccc	3300
atgacg	agg	gtt	ca	gtt	ca	gag	3360
ctgggtc	cc	gg	gg	gtt	gtt	gtt	3420
agacg	gg	cc	gg	gtt	gtt	gtt	3466
<212> Type	:	DNA					
<211> Length	:	3466					
SequenceName	:	SEQ ID 166					
SequenceDescription	:						

Final TCR Display PCT.workFile.txt

Custom Codon

Sequence Name : SEQ ID 166

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 KYLLPTAAAG LLLLAAQPAM AKQEVQTQIPA ALSVPEGENL VLNCSTDSA IYNLQWFRQD 60
 PGKGTLTSLLL IQSSSQREQTS GRLNASLDKS SGRSTLYIAA SQPGDSATYL CAVRPTSGGS 120
 YIPTFGRGTS LIVHPYIQNP DPAVYQLRDS KSSDKSVCLF TDFDSQTNVS QSKDSDVYIT 180
 DKCVLDMRSM DFKSNSAVAW SNKSDFACAN AFNNSIIPED TFFPSPESSM KKLLFAIPLV 240
 VPFYSHSAQA GVTQTPKFQV LKTGQSMTLQ CAQDMNHEYM SWYRQDPGMG LRLIHYSVGA 300
 GITDQGEVPN GYNVSRSTTE DFPLRLLSAA PSQTSVYFCA SSYVGNTGEL FFGEGSRLTV 360
 LEDLKNVFPP EVAVFEPSEA EISHTQKATL VCLATGFYPD HVELSWWVNG KEVHSGVCTD 420
 PQPLKEQPAL NDSRYALSSR LRVSATFWQD PRNHFRCQVQ FYGLSENDEW TQDRAKPVTQ 480
 IVSAEAWGRA DAAA 494
 <212> Type : PRT
 <211> Length : 494
 SequenceName : SEQ ID 167
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gatatccatgg ccataagtgg agtccctgtg ctaggatttt tcatcatagc tgtgctgatg 60
 agcgctcagg aatcatgggc tatcaaagaa gaacatgtga tcatccaggc cgagttctat 120
 ctgaatcctg accaattcagg cgagtttatg tttgactttg atggtgatga gatttccat 180
 gtggatatgg caaagaagga gacggctgg cggcttgaag aatttggacg atttgccagc 240
 tttgaggctc aaggtgcatt ggccaacata gctgtggaca aagccaacct ggaaatcatg 300
 acaaagcgct ccaactatac tccgatcacc aatgtacctc cagaggtAAC tttgctcacc 360
 aacagccctg tggaaactgag agagcccaac gtcctcatct gtttcatcga caagttcacc 420
 ccaccagtgg tcaatgtcac gtggcttcga aatggaaaac ctgtcaccac aggagtgtca 480
 gagacagtct tcctgcccag ggaagaccac cttttccgca agttccacta tctcccttc 540
 ctgccctcaa ctgaggacgt ttacgactgc agggtgagc actggggctt ggatgagcct 600
 cttctcaagc actggggagtt tggatgctcca agccctctcc cagagactac agagaacgtg 660
 gatgggggtc tgactgatac actccaagcg gagacagatc aacttgaaga caagaagtct 720
 gcgttgcaga ccgagattgc caatctactg aaagagaagg aaaaactaga gttcatcctg 780
 gcagcttacg gatctggtgg tggtctgaac gatattttt aagctcagaa aatcaatgg 840
 catgagtagg atcc 854
 <212> Type : DNA
 <211> Length : 854
 SequenceName : SEQ ID 168
 SequenceDescription :

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Custom Codon

Sequence Name : SEQ ID 168

Sequence

<213> OrganismName : Synthetic Sequence	
<400> PreSequenceString :	
aattctactc gtaaaagcgag ttgaaggatc atatttagtt gcgttatga gataagattg	60
aaagcacgtg taaaatgttt cccgcgcgtt ggcacaacta tttacaatgc ggccaagttt	120
taaaaagattc taatctgata tgaaaaaaa caccttgcg gcccgagttg tttgcgtacg	180
tgactagcga agaagatgtg tggaccgcag aacagatagt aaaacaaaaac cctagtattg	240
gagcaataat cgatttaacc aacacgtcta aatattatga tgggtgtcat tttttgcggg	300
cgggcctgtt atacaaaaaa attcaagtac ctggccagac tttgccgcct gaaagcatag	360
ttcaagaatt tattgacacg gtaaaaagaat ttacagaaaa gtgtcccgcc atgttggtgg	420
gcgtgcactg cacacacggt attaatcgca ccggttacat ggtgtgcaga tatttaatgc	480
acaccctggg tattgcgccg caggaagcca tagatagatt cggaaaaagcc agaggtcaca	540
aaattgaaag aaaaaattac gttcaagatttattaattatata tttgcattct	600
ttaacaataa ctttatccta tttcaaattt gttgcgccttc ttccagcgaa ccaaaactat	660
gcttcgcttg ctccgttttag ctgttagccg atcagtggcg ttgttccaat cgacggtagg	720
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gttttccacg tacgtctttt ggccggtaat agccgtaaac gtatgtccgt cgccgcgtcac	840
gcacaacacc ggatgttgc gcttgcgcgc ggggtattga accgcgcgat ccgacaaaatc	900
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cttttgcattt gtttcccttga agccgggtta catgcgggtt agatcagtca tgacgcgcgt	1020
gacctgcaaa tctttggcct cgatctgctt gtccttgatg gcaacgatgc gttcaataaa	1080
ctcttgggtt ttaacaagtt cctcgggtttt ttgcgccacc accgcttgca ggcgtttgt	1140
gtgctcggtg aatgtcgcaa tcagcttagt caccaactgt ttgctctcct cctccgttg	1200
tttgatcgcg ggatcgtaact tgccgggtgca gagcacttga ggaattactt cttctaaaag	1260
ccattcttgt aattctatgg cgtaaggcaa tttggacttc ataatcagct gaatcacgccc	1320
ggatttagta atgagcactg tatgcggctg caaatacagc gggcgcccc ttttcacgac	1380
gctgttagag gtggggcccc cattttggat ggtctgcata aataacgatt tgtattttt	1440
gtctacatga acacgtatag ctttatcaca aactgtatat tttaaactgt tagcgacgtc	1500
cttggccacg aaccggaccc gttggtcgcg ctctagcacg taccgcaggt tgaacgtatc	1560
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Final TCR Display PCT.WorkFile.txt
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 <211> Length : 10096
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 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 169

Sequence

<213> OrganismName : Synthetic Sequence
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 <212> Type : DNA
 <211> Length : 941
 SequenceName : SEQ ID 170
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 170

Sequence

<213> OrganismName : Synthetic Sequence
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<212> Type : PRT

Final TCR Display PCT.WorkFile.txt

<211> Length : 116
 SequenceName : SEQ ID 171
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
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<212> Type : PRT
 <211> Length : 116
 SequenceName : SEQ ID 172
 SequenceDescription :

Feature

Sequence: SEQ ID 172:
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 <222> LocationFrom : 1
 <222> LocationTo : 116
 Other Information : where x is e, q or r
 CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
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<212> Type : PRT
 <211> Length : 117
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 SequenceDescription :

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<213> OrganismName : Synthetic Sequence
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<212> Type : PRT
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<213> OrganismName : Synthetic Sequence
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<212> Type : PRT
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 SequenceDescription :

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